

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q2M2I8	AAK1_HUMAN	Homo sapiens	AP2-associated protein kinase 1	19.448366	T354;T359;T360;T441;S447;T448;T507;S519	S14;T234;S235;T354;T389;T441;T606;S618;T620;S623;S624;S637;S650;T653;T687;S731;S846;S937;S938	28657654;29351928;22564745;34019948;30620550	MKKFFDSRREQGSGSLGSGSSGGG GSTSGLGSGYIGRVFGIGRQQVTVDE VLAEGGFVAVFLVVRTSNGMKCALKR MFVNNNEHDLQVCKREIQIMRDLSSG HKNIVGYIDSSINNVSSGDVWEVLIL MDFCRGGQVVNLMNQRLQTGFTE NEVLQIFCDTCEAVARLHQCKTPIIH RDLKVENILLHDRGHYVLCDFGSAT NKFQNPQTEGVNAVEDEIKKYTTLS YRAPEMVNLYSGKIITTKADIWALGC LLYKLCYFTLPPFGESQVAICDGNFTI PDNSRYSQDMHCLIRYMLEPDPDK RPDIYQVSYFSFKLLKKECPINVQN SPIPAKLPEPVKASEAAAKKTQPKAR LTDPIPTTETSIAPRQRPKAGQTQPN PGILPIQPALTPRKRATVQPPQAAG SSNQPGLLASVPQPKPQAPPSPQLP QTQAKQPQAPPTPQQTTPSTQAQGLP AQAQATPQHQQQLFLKQQQQQQQP PPAQQQPAGTFYQQQQAQTQFQA VHPATQKPAIAQFPVVSQGGSQQL MQNFYQQQQQQQQQQQQQLATA LHQQQLMTQQAALQKQPTMAAGQQ PQPQAAAAPQAPAEPAIQAPVRQ QPKVQTTPPPQVQKQVGLTPPSSP KTQRAGHRRILSDVTHSAVFGVPAS KSTQLLQAAAAEASLNKSKSATTTPS GSPRTSQQNVYNPSEGSTWNPFD DNFSKLTAEELLNKDFAKLGEGKHP EKLGGSAESLIPGFQSTQGDFAFATS FSAGTAEKRKGGQTVDSGLPLLSVS DPFIPLQVPDAPEKLEGLKSPDTSLL LPDLLPMTDPFGSTSDAVIEKADVAV ESLIPGLEPPVPQRLPSQTESVTSNR TDSLTTGEDSLLDCSLLSNPTDLLEE FAPTAISAPVHKAEDSNLISGFDVP EGSDKVAEDEFDPIPVLITKNPQGGH SRNSSGSESSLPNLARSLLLVQQLI DL